

Sequence Listings

(1) GENERAL INFORMATION:

- 5 (i) INVENTORS: B. Mollet
R.D. Pridmore
M.C. Zwahlen
- 10 (ii) TITLE OF THE INVENTION: Novel strains of the *Bacillus subtilis* group
for food fermentation
- 15 (iii) NUMBER OF SEQUENCES: 4
- (iv) ADDRESS:
(A) Address: Winston & Strawn
(B) Street: 200 Park Avenue
(C) City: New York
(D) State: New York
(E) Country: USA
20 (F) Regional Code:
- 25 (v) COMPUTER READABLE FORM:
(A) Medium type: Floppy Disk
(B) Computer: IBM PC Compatible
(C) System: PC-DOS/MS-DOS
(D) Software: Patentin Release Nr. 1, Version Nr. 1,25
- 30 (vi) INFORMATION OF THE APPLICATION:
(A) Application number: to be obtained
(B) Application date: herewith
(C) Classification:
- 35 (vii) PRIORITY DATES: none
- (viii) Attorney/Agent Information
(A) Name: Allan A. Fanucci
(B) Registration number: 30,256
(C) Reference: 88265-4011
- 40 (ix) TELECOMMUNICATION:
(A) Telephone: 212-294-3311
(B) Telefax: 212-294-4700
- 45 (2) INFORMATION FOR SEQ ID. NO. 1:
(i) SEQUENCE CHARACTERISTICS:
(A) length: 30 nucleotides
(B) type: nucleic acid
(C) strandedness: single stranded
(D) topology: linear
- 50 (ii) MOLECULE TYPE: DNA

(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 1:
GCGGCGGATC CGCTGATGAT CTCCCAGCCC 30

5 (3) INFORMATION FOR SEQ ID. NO. 2:
(i) SEQUENCE CHARACTERISTICS:
(A) length: 44 nucleotides
(B) type: nucleic acid
(C) strandedness: single stranded
10 (D) topology: linear

(ii) MOLECULE TYPE: DNA

(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 2:
15 CTCAAATTCC ATTTCCCTCAT CAGGACATGC ATAGCGTATC ATCC 44

(4) INFORMATION FOR SEQ ID. NO. 3:
(i) SEQUENCE CHARACTERISTICS:
(A) length: 31 nucleotides
(B) type: nucleic acid
(C) strandedness: single stranded
20 (D) Topology: linear

25 (ii) MOLECULE TYPE: DNA

(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 3:
GGGGTCGAAT TCCACGAGAT ATCTAACTGC C 31

30 (5) INFORMATION FOR SEQ ID. NO. 4:
(i) SEQUENCE CHARACTERISTICS:
(A) length: 44 nucleotides
(B) type: nucleic acid
35 (C) strandedness: single stranded
(D) Topology: linear

(ii) MOLECULE TYPE: DNA

40 (iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 4:
GGATGATAACG CTATGCATGT CCTGATGAGG AAATGGAATTGAG 44